

FIGURE 1

Inventors: Volker SCHELLENBERGER, et al. Filing Date: October 10, 2001 Contact: David W. Maher Contact #: (650)849-4908 Attorney Docker No: 23623-7060 Sheet 2 of 2 FIGURE 2

						Section 1
	(1)	1 ,10	20	30	40	52
BL A sobria CAA56561 1	(1)	MKQTRALPLLAL	TLLLAPL	SLAAPVDPLK	AVVDDAIR	PVLKQHRIP
BL E cloacae P05364	(1)	-MMRKSLCCALLLG-	ISCSAL	ATPVSEKQLA	EVVANTIT	PLMKAQSVP
BL E coli CAB36900 1 BL O anthropi CAC17622 1	(1)	-MMKKSLCCALLLT- MRTSTTLLIGFLTT	AUTPNNGALA	ASKVNDGDLE	BIVNETLE	PLMQEQAIP
BL P aerocinosa S13408	/41	MRDTREPCTCGTAAS	STLLFATTPAL.	AGEAPADRLE	ALVIDA AUO	STUREMAND
BL S enteriditis CAA76196 1	741	MKKSLSATLISA-	LLAPSA	PGFSAADNVA	AVVDSTTY	DT.MAGG BTP
BL Y enterolitica P45460 Consensus	(1)	-MMKKSLINTLIFTS	SIATEPLITLA	QTALTELQVA A E OLA	IVD TI	PLEKQGIP
Consensus	(1)	MARSH MA				Section 2
	(53)	53 60	70	80	90	104
BL A sobria CAA56561 1	(48)	GMAVAVLKGGQAHY	FNYGLADVATG	akvneqtlee	IGSVSKTY	TATLGAYAV
BL E cloacae P05364 BL E coil CAB36900 1	(47)	GMAVAVIYQGKPHY:	YTFGKADIAAN FTEGKADIAAN	KPVTPQTLFE HPVTCOTLFE	LGSISKTF	TGVLGGDAI
BL O anthropi CAC17822 1	/E31	GMAVAITIDGKSHFI	EGYGVASKESG	QKVTEDTIFE	IGSVSKTF	TAMLEGYEL
RL P aeroginosa S13408	(F3)	GLAVAISLKGEPHY	FSYGLASKEDG	RRVTPETLEE	IGSVSKTF'	TATLAGYAL
BL S enteriditis CAA76196 1 BL Y enterolitica P45460	/52\	GMAVAVSVKGKPYY GMAVAVFYDGKPQF	FNYGMADIKAG	RPVTENTLE	LGSVSKTF	TGVLGAVSV
Consensus	(53)	GMAVAV GKPHY	F YGLADI AG	PVTEQTLEE	LGSVSKTF	TGVLGGYAI
		400 440	400	130	440	Section 3
	(105)	VKGGFKLDDQVSGH	120	THUSDIANU	140	156
DI E coll CAR36900 4	(QQ)	ARGETELSDPVTKY	WPELTGKOWOG	IRLLHLATYT	AGGLPLOI	PD DVRD~KA
BL O anthropi CAC17622 1 BL P aeroginosa S13408						
BL S enteriditis CAA76196 1 BL Y enterolitica P45460	(98)	AKKEMALNDPAAKY	QPELALPOWKG	ITLLDLATYT	AGGLPLOV	PDAVKS-RA
BL Y enterolitica P45460	(104)	AKG M L DPVTKY	APELTGSQWKD	VKMLHLATYT TTMLDLATYT	AGGLPLOL	PDSVTD-QK
Consensus	(105)	WEG W P DEALTI	EFF GDG# G		MAGRERA	Section 4
	(157)	157 1	70 ,18	0 19		208
BL A sobria CAA56561 1			AGTOROYSNES	IGLFGHLAAS	SLOQPEST	LMEQTLLPA
BL E coil CAB36900 1 BL O anthropi CAC17622 1	(156)	SMLAYFKNWKPDYP	AGTORRYSHPS	IGLFGYLAAF	SMOKPEDV	LMECKLLPA
BL P aeroginosa S13408	(157)	QIRDYYRQWQPTYA	PGSQRLYSNPS	IGLFGYLAAF TGLFGALTAN	ISLGQPFER	LMEQQVFPA
BL P aerodinosa S13408 BL S enteriditis CAA76196 1 BL Y enterolitica P45460	(149)	SIWQYYQQWQPQWA	PGVMRNYSNAS	IGLEGALAVE	REGLIFEN	AWKEAALds
Consensus	(157)	SLL YYQNWQP W	PGT RLYSNAS	IGLEG LAAR	SSGMPFE	LM RVL P
		209 220	230 MARYAFGYSKE	240 DKPIRVNPGI	25	Section 5 0 260 KTGSADLLA
BL A sobria CAA56561 1	(209) (203)	209 220	230 MARYAFGYSKE	240 DKPIRVNPGI	25 LADRAYGI	Section 5 0 260 KTGSADLLA
Bt. A sobria CAA56561 1 Bt. E cloacae P05364 Bt. E coil CAB36900 1	(209) (203) (202) (202)	209 220 LGLEHTYLQVPEAA LKLDETWINVPKAE LKLAHTWITVPQNE	230 MARYAFGYSKE EANYAWGYRDG QKDYAWGYREG	240 DKPIRVNPGI -KAVRVSPGI -XPVHVSPG	25 MLADEAYGI MLDAQAYGV MLDASAYGV	Section 5 0 260 KTGSADLLA KTNVQDMAN KSSVIDMAR
Bt. A sobria CAA56561 1 Bt. E cloacae P05364 Bt. E coil CAB36900 1	(209) (203) (202) (202)	209 220 LGLEHTYLQVPEAA LKLDETWINVPKAE LKLAHTWITVPQNE	230 MARYAFGYSKE EANYAWGYRDG QKDYAWGYREG	240 DKPIRVNPGI -KAVRVSPGI -XPVHVSPG	25 MLADEAYGI MLDAQAYGV MLDASAYGV	Section 5 0 260 KTGSADLLA KTNVQDMAN KSSVIDMAR
BL A sobria CAA56561 1 BL E cloacae P05384 BL E coil CAB38900 1 BL O anthroot CAC17622 1 BL P aerocinosa S13408	(209) (203) (202) (202) (208) (209)	209 220 LGLEHTYLQVPEAA LKLDETWINVPKAE LKLAETWITVPQNE FGLKNTFINVPESQ LGLEQTHLDVPEAA	230 MARYAFGYSKE EAHTAWGYRDG QKDYAWGYREG LHKNYAYGYSKA LAQYAQGYGK	240 DKPIRVNPGI -KAVRVS PGI -KPVHVS PGI DKPIRVS PGI -KPVRVS PGI	25 (LADBAYGI (LDAQAYGV (LDABAYGV ALDAQAYGI PLDABGYGV (LDABSYGV	Section 5 0 260 KTGSADLLA KTNVQDMAN KSSVIDMAR KKTTALDLAR KKTSAADLLR KSASKDMLR
Bt. A sobria CAA56561 1 Bt. E cloacae P05384 Bt. E coil CAB36900 1 Bt. O anthrool CAC17622 1 Bt. P aerocinosa S13408 Bt. S enteriditis CAA76196 1	(209) (203) (202) (202) (208) (209) (201)	209 220 LGLEHTYLQVPEAA LKLAETWINVPKAE LKLAETWITVPQNE FGLKNIFINVPEAA LGLEQTHLDVPEAA LGLSHTFITVPESA	230 MARYAFGYSKE EAHTAWGYREG QKDYAWGYREG MKNYAYGYSKA LAQYAQGYGK QSQYAYGYKNE QSQYAYGYKNE	240 DKPIRVNPGI -KAVRVSPGG -KPVHVSPGGI DRPLRVGPGI -RPVRVSPGG	25 MLADEAYGI MLDAQAYGV MLDAQAYGI MLDAQAYGI MLDAEGYGV MLGEEAYGV MLGEEAYGV	Section 5 0 260 KTGSADLLA KTHVQDMAN KSSVIDMAR KTTALDLAR KTTALDLAR KTSAADLLR KSASKDMLR
Bt. A sobria CAA56561 1 Bt. E cloacae P05384 Bt. E coil CAB36900 1 Bt. O anthrool CAC17622 1 Bt. P aerocinosa S13408 Bt. S enteriditis CAA76196 1	(209) (203) (202) (202) (208) (209) (201) (207) (209)	209 220 LGLEHTYLQVPEAA LKLDHEWINVPKAE LKLAHTWITVPQNE FGLKNIFINVPESQ LGLEQTHLDVPEAA LGLSHFITVPESA LKLDHTFITIPESM LGLDHTFI VPEA	230 MARYAFGYSKE EAHYAWGYRDG QKDYAWGYREG MKNYAYGYSKA LAQYAQGYGKO QSQYAYGYKNW QSNYAWGYKDG ANYAWGYK	240 DKPIRVNPG: -KAVRVSPG: -KPVHVSPG: MXPIRVSGGI DRPLRVGPG: -KPVRVSPG: KPVRVSPG	25 ALADBAYGI ALDAQAYGV ALDAQAYGI ALDAGAYGI ALDABGYGV ALDABGYGV ALGEBAYGV ALGEBAYGV ALGABAYGV	Section 5 0 260 KTGSADLLA KTNVQDMAN KSSVIDMAR KTTALDLAR KTSAADLLR KSSASKDMLR KSTSQDMVR KSTSS DMLR Section 6
BL A sobria CAA56581 1 BL E cloacae PG5384 BL E coil CAB36900 1 BL O anthrool CAG17622 1 BL P aerocincos 313406 BL S entercitiis CAA76196 1 BL Y entercitiica P45400 Consensus	(209) (203) (202) (202) (208) (209) (201) (207)	209 220 IGLEHTYLQVPBAA LKLDETWINVPKAE LKLAETWITVPGNE FGLKNITINVBSO LGLECTHLDVPBAA LGLSHTPILTVPBSA LKLDHTFILTVPBSA LKLDHTFILTVPBSA LGLDHTFILTVPBSA LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LGLDHTFILTPBSM LG	230 MARYAFGYSKE EAHIANGYRDG GKDYAFGYREG MKNYAYGYSKA LAQYAQGYGRU QSQYAYGYKNB QSNYAWGYKDG ANYAWGYK	240 DKPIRVNPGI -KAVRVSPGI -KPVHVSPGI -KPVHVSPGI -KPVRVSPGI -KPVRVSPGI -KPVRVSPGI -KPVRVSPGI	25 MLADERYGI MLDAQAYGV MLDAEAYGV ALDAEAYGV PLDAEGYGV MLGEEAYGV LDAEAYGV LDAEAYGV	Section 5 0 260 KTGSADLLA KTNVQDMAN KSSVIDMAR KTTALDLAR KTSADLLR KSASKDMLR KSASKDMLR KSASKDMLR KSTSQDMVR VKTSS DMLR Section 6 312
BL A sobria CAA56561 1 BL E cloacae P05394 BL E cell CAB5900 1 BL P acciding 813406 BL S enteriditis CAA76196 1 BL Y enterolitic AP4540 Consensus BL A sobria CAA56561 1	(209) (203) (202) (202) (208) (209) (201) (207) (209) (261) (255)	209 220 LGLEHTYLQVPBAA LKLDETWINVPKAE LKLAESWITVPCNE FGLKNIFINVPBSA LGLEGTHIDVPBAA LGLSHTPITVPBSA LKLDHTFITVPBSA	230 MARYAFGYSKE EAHYAWGYRDG MKNYAYGYSKA LAQYAQGYGKD QSQYAYGYKNB QSNYAWGYKDG ANYAWGYK	240 DKPIRVNPGIKAVRVSPGIKPURVSPGI DRPLRVSPGIKPURVSPGIKPURVSPGI	25 MLADEAYGI MLDAQAYGU ILDAEAYGU ALDAQAYGI PLDAESYGU ILDAESYGU ILDAEAYGU ILDAEAYGU ILDAEAYGU MSQGLGWES	Section 5 0 260 KTGSADLLA KTNVQDMAN KTSAVIDMAR KTTALDLAR KTSAADLLR KSASKDMLR KSASKDMLR KSISQDMVR KSISQDMVR SECTION 6 312
BL A sobria CAA56561 1 BL E cloacae P05394 BL E cell CAB5900 1 BL P acciding 813406 BL S enteriditis CAA76196 1 BL Y enterolitic AP4540 Consensus BL A sobria CAA56561 1	(209) (203) (202) (202) (208) (209) (201) (207) (209) (261) (255)	209 220 LGLEHTYLQVPBAA LKLDETWINVPKAE LKLAESWITVPCNE FGLKNIFINVPBSA LGLEGTHIDVPBAA LGLSHTPITVPBSA LKLDHTFITVPBSA	230 MARYAFGYSKE EAHYAWGYRDG MKNYAYGYSKA LAQYAQGYGKD QSQYAYGYKNB QSNYAWGYKDG ANYAWGYK	240 DKPIRVNPGIKAVRVSPGIKPURVSPGI DRPLRVSPGIKPURVSPGIKPURVSPGI	25 MLADEAYGI MLDAQAYGU ILDAEAYGU ALDAQAYGI PLDAESYGU ILDAESYGU ILDAEAYGU ILDAEAYGU ILDAEAYGU MSQGLGWES	Section 5 0 260 KTGSADLLA KTNVQDMAN KTSAVIDMAR KTTALDLAR KTSAADLLR KSASKDMLR KSASKDMLR KSISQDMVR KSISQDMVR SECTION 6 312
BL A sobria CAA56561 1 BL E cili CAA56561 1 BL E cili CAA56560 1 BL D aerocinosa \$134.06 BL S enteriditis CAA76196 1 BL Y enterditis CAA76196 1 BL Y enterditis CAA76196 1 BL A sobria CAA6561 1 BL E cili CAA6568 1 BL E cili CAB5680 1 BL B COII CAB5680 1 BL B COII CAB5680 1 BL B COII CAB5680 1 BL OBERGE POSS68	(209) (203) (202) (202) (208) (209) (201) (207) (209) (261) (255) (253) (263)	209 220 LGLEHTYLGYPEAR LKLARTWITVPGME LKLARTWITVPGME LGLEGTHLDVPEAR LKLARTWITVPEAR LKLDHTFIITVPEAR LKLDHTFIITVPEAR EVMANMPRENVADJ RVGANMDASEVGET RVGANMDASEVGET RVGLKIDSSSLE	230 MARYAFGYSKE EANTAMGYNG GKNYANGYNG MKNYAGYSK MKNYAGYSK GSQYAYGYKNH GSNYAMGYK 280 ANYAMGYK CELQGIALI LELKGGFAL LELKGGFAL LELKGGFAL	240 DKPIRVNPGIKAVHVSPGIKEVHVSPGI NKPIRVSGGI DRPIRVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEVHVSPGIKEV	25 MIADEAYGI MIDAQAYGV ALDAQAYGV LDAEGYGV LDAEGYGV MIGEEAYGV LDAEAYGV AGGEGAYGV MYGGLGWES MYGGLGWES MYGGLGWES MYGGLGWES	Section 5 0 260 KTGSADLLA KTNVQDMAN KSSVIDMAR KKTALDLAR KKTSADLLR KSASKDMLR KSTSQDMVR KSTSGDMVR Section 6 312 SYAYPYSEGT LINWPURANT LINWPURANT
BL A sobria CAA56561 1 BL E cloacae P05394 BL E coll CAA59500 1 BL S enterfold CAC17313408 BL S enterfold CAC17313408 BL S enterfold CAC17313408 BL Y enterollic CAA731341 BL C cloacae P05364 BL E cloacae P05364 BL E coll CAB38900 1 BL O anthroal CAC17822 1 BL P enterollicae 313408	(209) (203) (202) (202) (208) (209) (201) (207) (209) (261) (253) (253) (261) (261)	209 220 LGLERTYLOVPEAA LKLDERWINVPKAE LKLAETWITVPGWE FOLKNITINVPESO LGLEGITILOVPEAA LKLDHTFITVPESS LKLDHTFITVPESS LKLDHTFITVPESS LKLDHTFITVPESS LWANNAMABRUVADAW LWANNAMABRUVADAW LWANNAMABRUVADAW LWANNAMABRUVADAW LYGANNABASHOUEP LYGANIS - SELE LYG	230 MARYAFGYSKE ANTAMOSTROG GROYAMGYSKE GROYAMGYSKE GSGYAYGYSKE ANYAMGYS 280 CK-ALQQAIALISLKGGIALLSLKGGIALL E-BYAGALDAI R-PWAGALDAIDEMMAYA	240 DKPIRVNPGIKAVRVSPGIKPVRVSPGI ORPLRVPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPVRVSPGIKPV	25 MLADEA YGI MLDAQA YGY MLDAEA YGY MLDAEA YGY MLDAEA YGY MLGEEA YGY MLGEEA YGY MLGEGA YGY MYGGLGWEN MYGGLGWEN MYGGLGWEN MYGGLGWEN MYGGLGWEN MYGGLGWEN	Section 5 0 260 KITESADLLA KKINVODMAN KKINVODMAN KKINADLA KKINADLA KKINADLA KKINADLA KKINADLA KKINADLA KKINADA KINADA SECTION
BL A sobria CAA56861 1 BL E cloacae P05384 BL E cloacae P05384 BL O anthrool CAC17822 1 BL P arecdiness \$134.08 BL S enterditis CAA78186 1 BL V enterditis P46460 Comensus BL A sobria CAA65861 1 BL E cloacae P05384 BL E coli CAB63800 1 BL O anthrool CAC17822 1 BL P arecdiness \$134.08	(209) (203) (202) (202) (202) (203) (201) (207) (207) (209) (255) (253) (253) (253) (253) (253) (253) (253) (253)	209 220 LGLEHTYLOYBEA LKLAHTKINYBKAE LKLAHTKINYBKAE LKLAHTKINYBKAE LKLAHTKINYBKAE LKLAHTKINYBKAE LGLECHLLOYBEA LGLECHLLOYBEA LGLECHLLOYBEA LGLECHLLOYBEA LGLECHLLOYBEA LGLECHLOYBEA LGLECHL	MARYAFOTSIK BAHTANOTRIO GKDYANOTRIO GKDYANOTRIO GKDYANOTRIO GKDYANOTRIO GASOANOTRIO ANYANOTRIO BANTANOTRIO GANTANOTRIO K-ALQGIALI L-DEGRAVAA- L-TLQGIALI L-DEGRAVAA- L-DEGRAVA	240 DKPIRVNPGI	25 MIADEA YGI MIDAQAYGU PLDAE AYGU ALDAQAYGU PLDAE GYGV LDAE AYGU LDAE AYGU LDAE AYGU MAGGE AYGU MAGGE GWES MYGGL GWES	Section 5 0 260 KTGSADLLA KTNVQDMAN KSSVIDMAR KTTALDLAR KTSADLLR KSASKDMLR KSTSQDMVR KSTS DMLR Section 6 312 SYAYPYSEGT LLNH PVEAL LLNH PVEAL LLNH PVEAL KYDWFSALKR KYDWFSALKR KYDWFSALKR KYDWFSALKR KYDWFSALKR
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